

XP-002090820

22-01-98  
Complete 8

Hsaj03147.R55u031

ID HSAJ03147 standard; DNA; HUM; 239566 BP.  
AC AJ003147;  
NI e1246029  
DT 22-JAN-1998 (Rel. 54, Created)  
DT 03-JUL-1998 (Rel. 56, Last updated, Version 2)  
DE Homo sapiens complete genomic sequence between D16S3070 and  
DE D16S3275, containing Familial Mediterranean Fever gene disease  
XX  
KW HUMNK4 gene; mareno gene; marenostrin; metalloproteinase; mmp20 gene;  
KW olfactory receptor; zinc finger protein; znfmf gene.  
OS Homo sapiens (human)  
OC Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates;  
OC Catarrhini; Hominidae; Homo.  
RN [1]  
RA Bernot A., Heilig R., Clepet C., Smaoui N., da Silva C., Petit J.L.,  
RA Devaud C., Chiannilkulchai N., Fizames C., Samson D., Cruaud C.,  
RA Caloustian C., Gyapay G., Delpech M., Weissenbach J.;  
RT "A transcriptional map of the FMF region";  
RL Genomics 50:147-160(1998).  
RN [2]  
RP 1-239566  
RA Bernot A.;  
RT ;  
RL Submitted (07-JAN-1998) to the EMBL/GenBank/DDBJ databases.  
RL GENOSCOPE - Centre National de Sequencage, 2 rue Gaston Cremieux, EVRY  
RL BP191, FRANCE.  
DR SPTREMBL; O15361; O15361.  
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 214151 GGCTCATGCC TGTATTCCCG GCACTCTGGG AGGCCGAGGC GGGAGGATCA  
 214201 CCTGAGATCA GGAGTTCAAG ACCAACAGGG TGAAACCCCG TCTCTACTAA  
 214251 AAATACAAAA ATAAGCTGAG CGTGGTGGCG CATGCCTGTA ATCCCAGCTT  
 214301 CTCCAGAGGC CAAGGTGGGA GGATCATTTG AGGTCAGGAG TTCGAGACCA  
 214351 GCCTGGCCAA CATGGCAAAA CCCCCTTCCT ACTAAAAATA AGAAACTCAG  
 214401 CCAGACGCGG TGGTGCGCAC CTGTAGTCCC AGCTACTCGG GAGGCTGAGG

214451 CATGAGAATC GCTTGAGCCC GGGATGTGGA GGTGTCAGTG AGCTGAGATC  
214501 GGGCCACTGC ACTCCAGCCT GAGTGACAGA GCAAGACTCC ATCTCAAAAA  
214551 CAAAAGTTCA GGACTTCCTA GACTAGACAA TGGCAGTCAA TTACCATTCA  
214601 GTAAGAACAT GCTTGATGTT GCTATGTATT TCTCTCTCTC TTATTTTATT  
214651 TTTTGAGACA GGATCTTGCT CTGTTGCCCT GATGTGATCT CAGCTCACTG  
214701 CAACCTCTGC CTCCCAGGCT CAAGCAATCC TCTCACCTCA GCCTGTTTTT  
214751 TGTTTTTTCA ATAGCTGAAA ATACAGGTGT GTGCTACCAC GCCTGGCTCA  
214801 TTTTTTTGTA TTTTTGGTAG AGACAGGGTT TCACCATGTT GCCCAAGCTG  
214851 GTCTCAAACCT CCTGAGCTCT AGCAATCCAC CCACCTTAGC CTCCCCAAGT  
214901 GCTGAGATTA TTTATAAGTG TGAGCCACCA TGCCTAGCCT ATATTTCTCA  
214951 AATAGTACTT TTCAAACGTT TTAAGCAGGA CAACTGTTTC AAATAAGATC  
215001 TTACACAGGT CAGCAAAACC CAGATTATAC GCTTTAAAAA TAAAAATAAG  
215051 CCGGGCACGG TGGCCTATGT CTGAAATCCC AGCACTTTGG GAGGCTGAGG  
215101 CGGGAGGATC ACTTGAGGTC AGGAGTTCGA GACCAGCCTG GCCAACATGG  
215151 TGAACAGCAC TGAGAAGATT CTATACCAAC CCCGTTTATA TGATTGCATA  
215201 GCAATCCCTT TTTGCTAACC TAGAGATGTT TGTCTGACAT GAGTATTAAT  
215251 CATAAATGTA GTGAAGAAGT CTCAAAGAAC AGGTGTTCCA GCTCTGCCT  
215301 TTCGTGACCC TGACCTGCTT CTTAAAAAAC CCATTAGGAG CCTGAAGGAA  
215351 GTTTACTAAC CCAGTTCCAA AGGCCAGAGT GAAGAAAGGG ACGTTCCTGA  
215401 ACTAAAGTCA TCTGGATTTT GGTAGACCTG AGACTCCCAA TCCCCAGGTC  
215451 AGAGTGAGCT GCTCTGAGCT CCTGGTCCCC TTCCACAAA AGCAGCCAGC  
215501 ACTCAGCACT GGATGAGGAG GAGGCCTGGG CCCGCTTACC CTGAATGGCT  
215551 GCCCTGTGGA GCTCCTCGGC CAGCAGGCGC TGGTTGATGG CCCGCAGGAC  
215601 CTGCAGGGTG AGCTGCACGG CGTACTCTTC CCCATAGTAG GTGACCAGCA  
215651 GAGTGGCCAT CTTACCGGC CTGGCTCTCT GGATCTGGCT CCGGGGGATC  
215701 CTGGAGTGCT CCTTCTGCAC ACTGGTGTTC TGCAGCTTGA ACTTGAACCT  
215751 CTCGAAGTCA TAGGGCACCA GCTCCTCCAG GGTGGACAGC AGATGGTCAC  
215801 TAGGGGTCTT AGCCATGGTG CTGAGCAGGA GAGGCTCGAG CCAGCTGTCT  
215851 GGCTTCTGGT AGGAAAAGAA GCCTCTGTCC TTGGTGAGCA AGAAAAGGCA  
215901 GGTGTGAAA TAGCGAAAA GGCACAGGAA ATGCTCTGTG TCTTGGTGGG